STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information
Center (STIC) detected errors when processing the following computer readable
form:

Application Serial Number: _10/569,330_
Source: _IRJP_
Date Processed by STIC: _12/29/06_

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:
1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE
   APPLICANT, WITH A NOTICE TO COMPLY or,
2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A
   NOTICE TO COMPLY .
FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT
MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER
VERSION 4.4.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND
TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:
   http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is
a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.
Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the
United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm> , EFS Submission
   User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):
   U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street,
   Alexandria, VA 22314

Revised 01/10/06

Raw Sequence Listing Error Summary

ERROR DETECTED   SUGGESTED CORRECTION   SERIAL NUMBER: *10/569,330*

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1____Wrapped Nucleics   The number/text at the end of each line "wrapped" down to the next line. This may occur if your file
     Wrapped Aminos   was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will
                        prevent "wrapping."

2____Invalid Line Length   The rules require that a line not exceed 72 characters in length. This includes white spaces.

3____Misaligned Amino   The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers;
     Numbering   use space characters, instead.

4____Non-ASCII   The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please
                        ensure your subsequent submission is saved in ASCII text.

5____Variable Length   Sequence(s)____contain n's or Xaa's representing more than one residue. Per Sequence Rules,
                        each n or Xaa can only represent a single residue. Please present the maximum number of each
                        residue having variable length and indicate in the <220>-<223> section that some may be missing.

6____PatentIn 2.0   A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid
     "bug"   sequences(s) _____. Normally, PatentIn would automatically generate this section from the
                        previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to
                        the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for
                        Artificial or Unknown sequences.

7____Skipped Sequences Sequence(s)_____ missing. If intentional, please insert the following lines for each skipped sequence:
     (OLD RULES)   (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
                        (i)   SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
                        (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
                        This sequence is intentionally skipped
                        Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.

8____Skipped Sequences Sequence(s)_____ missing. If intentional, please insert the following lines for each skipped sequence.
     (NEW RULES)   <210> sequence id number
                        <400> sequence id number
                        000

9____Use of n's or Xaa's   Use of n's and/or Xaa's have been detected in the Sequence Listing.
     (NEW RULES)   Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
                        In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

10___Invalid <213>   Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or
     Response   scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or
                        is Artificial Sequence. (see item 11 below)

11___Use of <220>   Sequence(s)_____ missing the <220> "Feature" and associated numeric identifiers and responses. Use
                        of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown."
                        Please explain source of genetic material in <220> to <223> section or use "chemically synthesized" as
                        explanation. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32), also Sec. 1.823 of
                        Sequence Rules.

12___PatentIn 2.0   Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file,
     "bug"   resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence
                        listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

13___Misuse of n/Xaa   "n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid

IFWP

RAW SEQUENCE LISTING   DATE: 12/29/2006
PATENT APPLICATION: US/10/569,330   TIME: 15:38:03

Input Set : A:\L7350.0006 SEQUENCE LISTING.TXT
Output Set: N:\CRF4\12292006\J569330.raw

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 3 <110> APPLICANT: NAKAJIMA, Toshihiro
 4        AMANO, Tetsuya
 5        TSUCHIMOCHI, Kaneyuki
 6        YAMAZAKI, Satoshi
 7        YAGISHITA, Naoko
 9 <120> TITLE OF INVENTION: Synoviolin promoter
11 <130> FILE REFERENCE: L7350.0006
13 <140> CURRENT APPLICATION NUMBER: 10/569,330
14 <141> CURRENT FILING DATE: 2006-02-21
16 <150> PRIOR APPLICATION NUMBER: PCT/JP2004/012424
17 <151> PRIOR FILING DATE: 2004-08-23
19 <150> PRIOR APPLICATION NUMBER: JP2003-297913
20 <151> PRIOR FILING DATE: 2003-08-21
22 <160> NUMBER OF SEQ ID NOS: 13
24 <170> SOFTWARE: PatentIn version 3.3
26 <210> SEQ ID NO: 1
27 <211> LENGTH: 3046
28 <212> TYPE: DNA
29 <213> ORGANISM: Mus musculus
31 <400> SEQUENCE: 1
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36 gagtgctggg attaaaggta ggcgccacca cgcccagctt tttttttttt agataggatc     180
38 tcactctata gctgtacgct ggcctcagat ttatgatgct cttcctgcct cagtctccca     240
40 attttctggg attgtaggag tgggccacta tgctctgctc actacatgat ttcagaggtt     300
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52 ttaactgtta aagacaacag gtggtggtga agatggctga gaccaagagc acagggctga     660
54 ggggcagaca ggcactgaca ctgctaccct ttaatacagt tcctcctgtt gtgatcccca     720
56 accataatta cttcgttgct acttcataac tgtaattttg ctagttatga attgtaagta     780
58 aacgtctgat atgcaggata tctcatttgt gacccctgtg taacggtttg attcccaaag     840
60 ggcttacgac tcacaggttg agagccagcc actgccttaa agtcgtctag aatcagtttt     900
62 ctttcttttt tgacagacaa gatgtttaat tccgttgtac tgaaggaaag ccattttatg     960
64 tattttttct aagtgctcta tcagtaatga caattctgaa agcccctgtg ttatatttta    1020
66 acaacacagt cacctccggt tctgtattca ctgtccgtgt tgtgactccc acagtataaa    1080
68 ttcctccagt tgatcttcat gaattcttat atttgatccc ccccccctt aggcctctga    1140
70 attccgagtg agtccgattt aaaaatggga ggagcaccct ctagctgata aacctgggta    1200
72 atgaggtgtc cgctttcagt ttccattctg tacgcgacta tactgcttgt gtgagcccta    1260
74 acagacagaa tcagctcaga acaaagggtc tggctatctc ccaggatga acacgcacgc    1320
76 cgactgagct tttggggtgt tgaaaagtca acgccttcgc acagaactct ccaccccaac    1380
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Output Set: N:\CRF4\12292006\J569330.raw

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 82 ctcagctggc atccagctgc cttggcaccc agtccggggc cactctgcct acagaccdta   1560
 84 gcaaccactc acctgctttt ctttccctat aggccagaaa tttttccttt cttttctcat   1620
 86 tggtccgcgt aactttatcg caaccaatcg gcggtacacg ggaacaaact cactcctaca   1680
 88 caacctgcgt tgggggggagg taacctggga agacctatat ctgtttctg caccgctatt   1740
 90 tttttccgag aagcacttaa cttcttaccg tgtcgtagct atccctggaa tgaggcgctt   1800
 92 acacatttta tttctttcat gcctgacata aagtctggcc cttgctcgct cctgcccccc   1860
 94 gtccaaatgg ctcggcccgc ggaacgccca tcttccaggc acattgagag ccggagtctt   1920
 96 ggagggagtt tagggtggtg attctacaac ggcgactagc aagtggcggg cttcagccct   1980
 98 ttcccgctgc tctcctggtc gcgaccacac gtcacagctc tcgctcgttc cggttgctcg   2040
100 cgcaggggt ggggagtgtt gttaaccgga gcggctgccg cagtcgcggt gattgagcgt   2100
102 actccgccgc gccccgcgcc gccggaagtg aggtgtctta cccccgaagt tccggttcgc   2160
104 aggggtggg gagtgttgtt aaccggagcg gctgccgcag tcgcggtgat tgagcgtgct   2220
106 cgcggcgctg ggctcctggt gagtgggcct ggtcctgatt ggggttgggg ggtcggcgtc   2280
108 taggaccttg tcctttgggg tcactgcgat cagcccgccc cgctgcgttc ggccgccagt   2340
110 tttcggcctg tcagatggct ggagacctta ggcggcggcg cggccaccgt tccagaggcc   2400
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114 ctgactggtt ctcagccccg ccgccgcacc ctggcggtcg gccgtttctc cggttctcag   2520
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118 cgcgcgtctg cgctgaccac cctggcacag ctgtcactgg ttgtgtcgcc ttctcaagct   2640
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124 tcagtgactc cctagaggag gcgccttggc agacagcgtg gaagagccct agatttgaaa   2820
126 cgagattgat ccaagttcta ggccttgcat cagtgtgagc ctctaacccc tttgagtcct   2880
128 agtttctcgt ttgtgaaaca gggagtatat gctgttttga atctaatggc tgtcaaggtg   2940
130 aaatgagtgt ttgcccttac actctgccag ggactgtgct aggtttacat agtgtggata   3000
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138 <213> ORGANISM: Homo sapiens
140 <400> SEQUENCE: 2
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145 cctttattct gttctacttt tttcctatag cactgatcat cttccagcgt attagatttt   180
147 tcacttatgt ctgtggtttg ctgtcacatc tactaggata agctccacaa aggtagagat   240
149 ctttattttg ttcactgaca tcctaagtcc ctagaacagg agacacttga tccatatttg   300
151 tagactaact gaataaatga cttaattacc agtttggatg tggggcaga tagtgagcat   360
153 gatgcccgtt tccggagctg gggtgcagac agtgtctagg gacactgaac tgttttaaaa   420
155 gcaggataga tcccggctgg agaccacaca aggaaatcat cagcacctgg gtcaggggct   480
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159 gaggacacat gcatagccag agagaaaaga ggagcagagg catgtggatc acagaagctt   600
161 agggaggaga ctttcaagaa ggggagagag gttgagtcaa gcaagggctg aaagccaacc   660
163 attggatgca gtcactagaa agttacagat aggcaaggtg ttgtggctca cgcctgtaat   720
165 cccaacacct tgtggggctg aggtgggagg atcgcttgag cccgggaggt cgaggctgca   780
167 atgagccctg atggcgccaa tgcactccag cctgggcgac agagcaagac cctgtcgcaa   840
169 aaattaataa ataaataaat aaaagaaaa ggggaaaaa aagttatacg tggccttacg   900
171 gggaagccaa ctctgactgg ttataagctg aaactgtcaa gtcaacaggt ggcagggaag   960
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RAW SEQUENCE LISTING   DATE: 12/29/2006
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175 caggttttgg taagcctttg aatttcaatt gccccacgtt tcggggagg gggtagcacc 1080
177 ccctagctca taaaccttag tgattgatga ttaaatgaga tgacggagga aaacgcaagg 1140
179 cacaaagtgg atgcattagc tccattttgt taatcagcag gcttagttgg ctgcgaccca 1200
181 gacacgaact aaaatacagt gcagcccagg accagtgggg gtcttgctta tggctcagag 1260
183 ctgaacaaca catgggcagc aaaatcagac actgagatgc gggcaggcct gcgacgctga 1320
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187 cagtctcggg tacacctggt tgtccctccc tgtcctggcg cggcaaacgt tcccggaggc 1440
189 cagccaggga tcactcgccc aaggactgag ctttccctac tctcagccaa ctggagcggg 1500
191 accagggcct aggcaacgca gctgtccgcc cctaacaacc actcacctgc tttccccttt 1560
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211 cccacgccgc gccccgcgcc gccggaagtg aggtgtctga cccccgaagt tccggttcgc 2160
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217 gacccgaacc ttccccttga ggtctccgga gtcggcacgc ccctcagccc cgccgcacgc 2340
219 tttcggcctg tcagctggcc ggagacctca gacgccggtg cggccgcttt gctcaagcct 2400
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231 cctggagcag atttctgctg ttgcctccca ccccatcttc tccaccggag ggtcagcggt 2760
233 gcagctcccc ctcctccaac attgcagctt ttcctcatca cctccctaga ggaggcggct 2820
235 tggcaggcag cgtggaaaga gccctagatt tgaagcaaga ctgacccagg ttccaggcct 2880
237 tgcgtcagtg tgatcactta accccttcga gtctaatttg taaaatgggg tagcgtaagc 2940
239 tattctttgt ctgatgattt cgaggggcgaa atgtgatttc ccccccactt tctcctatga 3000
241 attgaggctg tgccaggcac cgggctattt tgcacagcac gagcatcaca taagttattt 3060
243 tcttgcccca tgcaggtctc cgggccaggg ca                                 3092
246 <210> SEQ ID NO: 3
247 <211> LENGTH: 19
248 <212> TYPE: DNA
249 <213> ORGANISM: Artificial
251 <220> FEATURE:
252 <223> OTHER INFORMATION: synthetic DNA
254 <400> SEQUENCE: 3
255 gcgccgccgt aagtgaggt                                               19
258 <210> SEQ ID NO: 4
259 <211> LENGTH: 20
260 <212> TYPE: DNA
261 <213> ORGANISM: Artificial
263 <220> FEATURE:
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Output Set: N:\CRF4\12292006\J569330.raw

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271 <211> LENGTH: 20
272 <212> TYPE: DNA
273 <213> ORGANISM: Artificial
275 <220> FEATURE:
276 <223> OTHER INFORMATION: synthetic DNA
278 <400> SEQUENCE: 5
279 actccgccaa gccccgcgcc                                         20
282 <210> SEQ ID NO: 6
283 <211> LENGTH: 16
284 <212> TYPE: DNA
285 <213> ORGANISM: Artificial
287 <220> FEATURE:
288 <223> OTHER INFORMATION: synthetic DNA
290 <400> SEQUENCE: 6
291 gcgccgccgg aagtga                                             16
294 <210> SEQ ID NO: 7
295 <211> LENGTH: 16
296 <212> TYPE: DNA
297 <213> ORGANISM: Artificial
299 <220> FEATURE:
300 <223> OTHER INFORMATION: synthetic DNA
302 <400> SEQUENCE: 7
303 gcgccgccgt aagtga                                             16
306 <210> SEQ ID NO: 8
307 <211> LENGTH: 101
308 <212> TYPE: DNA
309 <213> ORGANISM: Homo sapiens
311 <400> SEQUENCE: 8
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314 caggggtgg ggagtgttgt taaccggagg ggcagccgca g                       101
317 <210> SEQ ID NO: 9
318 <211> LENGTH: 101        musculus
319 <212> TYPE: DNA
320 <213> ORGANISM: Mus (musclus)
322 <400> SEQUENCE: 9
323 actccgccgc gccccgcgcc gccggaagtg aggtgtctct accccccgaag ttccggttcg  60
325 caggggtgg ggagtgttgt taaccggagc ggctgccgca g                       101
328 <210> SEQ ID NO: 10
329 <211> LENGTH: 11
330 <212> TYPE: DNA
331 <213> ORGANISM: Homo sapiens
333 <400> SEQUENCE: 10
334 gccggaagtg a                                                  11
337 <210> SEQ ID NO: 11
338 <211> LENGTH: 11
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RAW SEQUENCE LISTING   DATE: 12/29/2006
PATENT APPLICATION: US/10/569,330   TIME: 15:38:03

Input Set : A:\L7350.0006 SEQUENCE LISTING.TXT
Output Set: N:\CRF4\12292006\J569330.raw

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340 <213> ORGANISM: Artificial
342 <220> FEATURE:
343 <223> OTHER INFORMATION: mutant
345 <400> SEQUENCE: 11
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349 <210> SEQ ID NO: 12
350 <211> LENGTH: 10
351 <212> TYPE: DNA
352 <213> ORGANISM: Homo sapiens
354 <400> SEQUENCE: 12
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361 <213> ORGANISM: Artificial
363 <220> FEATURE:
364 <223> OTHER INFORMATION: mutant
366 <400> SEQUENCE: 13
367 gccaagcccc                                                      10
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insufficient explanation (give source of genetic material) see item 11 on Error Summary Sheet

RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 12/29/2006
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Input Set : A:\L7350.0006 SEQUENCE LISTING.TXT
Output Set: N:\CRF4\12292006\J569330.raw

<u>Invalid <213> Response:</u>
Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:3,4,5,6,7,11,13

VERIFICATION SUMMARY   DATE: 12/29/2006
PATENT APPLICATION: US/10/569,330   TIME: 15:38:04

Input Set : A:\L7350.0006 SEQUENCE LISTING.TXT
Output Set: N:\CRF4\12292006\J569330.raw